



2590
2570

RAW SEQUENCE LISTING ERROR REPORT

The Biotechnology Systems Branch of the Scientific and Technical Information Center (STIC) detected errors when processing the following computer readable form:

Application Serial Number: 10/027,736
Source: OPE
Date Processed by STIC: 5/13/02

THE ATTACHED PRINTOUT EXPLAINS DETECTED ERRORS.

PLEASE FORWARD THIS INFORMATION TO THE APPLICANT BY EITHER:

- 1) INCLUDING A COPY OF THIS PRINTOUT IN YOUR NEXT COMMUNICATION TO THE APPLICANT, WITH A NOTICE TO COMPLY or,
- 2) TELEPHONING APPLICANT AND FAXING A COPY OF THIS PRINTOUT, WITH A NOTICE TO COMPLY

FOR CRF SUBMISSION QUESTIONS, PLEASE CONTACT MARK SPENCER, 703-308-4212.

FOR SEQUENCE RULES INTERPRETATION, PLEASE CONTACT ROBERT WAX, 703-308-4216.

PATENTIN 2.1 e-mail help: patin21help@uspto.gov or phone 703-306-4119 (R. Wax)

PATENTIN 3.0 e-mail help: patin3help@uspto.gov or phone 703-306-4119 (R. Wax)

TO REDUCE ERRORED SEQUENCE LISTINGS, PLEASE USE THE CHECKER
VERSION 3.1 PROGRAM, ACCESSIBLE THROUGH THE U.S. PATENT AND
TRADEMARK OFFICE WEBSITE. SEE BELOW FOR ADDRESS:

<http://www.uspto.gov/web/offices/pac/checker>

Applicants submitting genetic sequence information electronically on diskette or CD-Rom should be aware that there is a possibility that the disk/CD-Rom may have been affected by treatment given to all incoming mail.

Please consider using alternate methods of submission for the disk/CD-Rom or replacement disk/CD-Rom.

Any reply including a sequence listing in electronic form should NOT be sent to the 20231 zip code address for the United States Patent and Trademark Office, and instead should be sent via the following to the indicated addresses:

1. EFS-Bio (<<http://www.uspto.gov/ebc/efs/downloads/documents.htm>> , EFS Submission User Manual - ePAVE)
2. U.S. Postal Service: U.S. Patent and Trademark Office, Box Sequence, P.O. Box 2327, Arlington, VA 22202
3. Hand Carry directly to:
U.S. Patent and Trademark Office, Technology Center 1600, Reception Area, 7th Floor, Examiner Name, Sequence Information, Crystal Mall One, 1911 South Clark Street, Arlington, VA 22202
Or
U.S. Patent and Trademark Office, Box Sequence, Customer Window, Lobby, Room 1B03, Crystal Plaza Two, 2011 South Clark Place, Arlington, VA 22202
4. Federal Express, United Parcel Service, or other delivery service to: U.S. Patent and Trademark Office, Box Sequence, Room 1B03-Mailroom, Crystal Plaza Two, 2011 South Clark Place, Arlington, VA 22202

Raw Sequence Listing Error Summary

01PE

ERROR DETECTED

SUGGESTED CORRECTION

SERIAL NUMBER: 101027,736

ATTN: NEW RULES CASES: PLEASE DISREGARD ENGLISH "ALPHA" HEADERS, WHICH WERE INSERTED BY PTO SOFTWARE

- 1 Wrapped Nucleics
 Wrapped Aminos The number/text at the end of each line "wrapped" down to the next line. This may occur if your file was retrieved in a word processor after creating it. Please adjust your right margin to .3; this will prevent "wrapping."

- 2 Invalid Line Length The rules require that a line not exceed 72 characters in length. This includes white spaces.

- 3 Misaligned Amino
 Numbering The numbering under each 5th amino acid is misaligned. Do not use tab codes between numbers; use space characters, instead.

- 4 Non-ASCII The submitted file was not saved in ASCII(DOS) text, as required by the Sequence Rules. Please ensure your subsequent submission is saved in ASCII text.

- 5 Variable Length Sequence(s) contain n's or Xaa's representing more than one residue. Per Sequence Rules, each n or Xaa can only represent a single residue. Please present the maximum number of each residue having variable length and indicate in the <220>-<223> section that some may be missing.

- 6 PatentIn 2.0
 "bug" A "bug" in PatentIn version 2.0 has caused the <220>-<223> section to be missing from amino acid sequences(s) . Normally, PatentIn would automatically generate this section from the previously coded nucleic acid sequence. Please manually copy the relevant <220>-<223> section to the subsequent amino acid sequence. This applies to the mandatory <220>-<223> sections for Artificial or Unknown sequences.

- 7 Skipped Sequences
 (OLD RULES) Sequence(s) missing. If intentional, please insert the following lines for each skipped sequence:
 (2) INFORMATION FOR SEQ ID NO:X: (insert SEQ ID NO where "X" is shown)
 (i) SEQUENCE CHARACTERISTICS: (Do not insert any subheadings under this heading)
 (xi) SEQUENCE DESCRIPTION:SEQ ID NO:X: (insert SEQ ID NO where "X" is shown)
 This sequence is intentionally skipped

 Please also adjust the "(ii) NUMBER OF SEQUENCES:" response to include the skipped sequences.

- 8 Skipped Sequences
 (NEW RULES) Sequence(s) missing. If intentional, please insert the following lines for each skipped sequence.
 <210> sequence id number
 <400> sequence id number
 000

- 9 ✓ Use of n's or Xaa's
 (NEW RULES) Use of n's and/or Xaa's have been detected in the Sequence Listing.
 Per 1.823 of Sequence Rules, use of <220>-<223> is MANDATORY if n's or Xaa's are present.
 In <220> to <223> section, please explain location of n or Xaa, and which residue n or Xaa represents.

- 10 Invalid <213>
 Response Per 1.823 of Sequence Rules, the only valid <213> responses are: Unknown, Artificial Sequence, or scientific name (Genus/species). <220>-<223> section is required when <213> response is Unknown or is Artificial Sequence

- 11 Use of <220> Sequence(s) missing the <220> "Feature" and associated numeric identifiers and responses.
 Use of <220> to <223> is MANDATORY if <213> "Organism" response is "Artificial Sequence" or "Unknown." Please explain source of genetic material in <220> to <223> section.
 (See "Federal Register," 06/01/1998, Vol. 63, No. 104, pp. 29631-32) (Sec. 1.823 of Sequence Rules)

- 12 PatentIn 2.0
 "bug" Please do not use "Copy to Disk" function of PatentIn version 2.0. This causes a corrupted file, resulting in missing mandatory numeric identifiers and responses (as indicated on raw sequence listing). Instead, please use "File Manager" or any other manual means to copy file to floppy disk.

- 13 Misuse of n n can only be used to represent a single nucleotide in a nucleic acid sequence. N is not used to represent any value not specifically a nucleotide.



Does Not Comply
Corrected Diskette Needed

OIKE

Error on p. 3

RAW SEQUENCE LISTING
PATENT APPLICATION: US/10/027,736

DATE: 05/13/2002
TIME: 09:37:07

Input Set : A:\Sequence List.txt
Output Set: N:\CRF3\05102002\J027736.raw

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3 <110> APPLICANT: Presta, Leonard G.
4   Namenuk, Angela K.
6 <120> TITLE OF INVENTION: NON-HUMAN PRIMATE Fc RECEPTORS AND METHODS OF USE
8 <130> FILE REFERENCE: 11669.92US01
10 <140> CURRENT APPLICATION NUMBER: US 10/027,736
C--> 11 <141> CURRENT FILING DATE: 2002-04-25
13 <160> NUMBER OF SEQ ID NOS: 72
15 <170> SOFTWARE: PatentIn version 3.1
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37 ggtgaataca ggtgccagag aggtccctca gggcgaagtg accccataca gctggaaatc      300
39 cacagagact ggctactact gcaggtatcc agcagagtct tcacagaagg agaacctctg      360
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49 ccgctcctgg aggggaatct ggtcaccttg agctgtgaaa caaagttgct tctgcagagg      660
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53 acgtcctctg aataccaaat actaactgct agaagagaag actctgggtt ttactggtgc      780
55 gaggccacca cagaagacgg aaatgtcctt aagcgcagcc ctgagttgga gcttcaagtg      840
57 cttggcctcc agttaccaac tcctgtctgg cttcatgtcc ttttctatct ggtagtgga      900
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61 aaaaagtgga atttagaaat atctttggat tctgctcatg agaagaaggt aacttccagc     1020
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69 <213> ORGANISM: Homo sapiens
71 <220> FEATURE:
72 <221> NAME/KEY: misc_feature
73 <222> LOCATION: (1)..(1128)
74 <223> OTHER INFORMATION: FcgammaRI alpha-chain

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82 ttgactgtg aggtgctoca tctgcctggg agcagctcta cacagtgggt tctcaatggc      180
84 acagccactc agacctcgac cccagctac agaatacct ctgccagtgt caatgacagt      240
86 ggtgaataca ggtgccagag aggtctctca gggcggaagt accccataca gctggaaatc      300
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90 gccttgaggt gtcatgcgtg gaaggataag ctggtgtaca atgtgcttta ctatcgaaat      420
92 ggcaaagcct ttaagttttt ccactggaat tctaacctca ccattctgaa aaccaacata      480
94 agtcacaatg gcacctacca ttgctcaggc atgggaaagc atcgctacac atcagcagga      540
96 atatctgtca ctgtgaaaga gctatttcca gctccagtgc tgaatgcac tgtgacatcc      600
98 ccactcctgg aggggaatct ggtcaccctg agctgtgaaa caaagttgct cttgcagagg      660
100 cctggtttgc agctttactt ctctttctac atgggcagca agaccctgcg aggcaggaac      720
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106 cttggcctcc agttaccaac tcctgtctgg ttctatgtcc ttttctatct ggcagtggga      900
108 ataatgtttt tagtgaacac tgttctctgg gtgacaatac gtaaagaact gaaaagaaag      960
110 aaaaagtggg atttagaaat ctctttggat tctggtcatg agaagaagg aatttccagc      1020
112 cttcaagaag acagacattt agaagaagag ctgaaatgtc aggaacaaaa agaagaacag      1080
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120 <213> ORGANISM: Cynomolgus

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125 <223> OTHER INFORMATION: FcgammaRIIA

128 <400> SEQUENCE: 3

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133 cctggatcca acgtgctccg ggaggactct gtgactctga cgtgcggggg cgctcacagc      180
135 cctgacagcg actccactca gtggttccac aatgggaatc gcacccccac ccacacacag      240
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163 <211> LENGTH: 936

164 <212> TYPE: DNA

165 <213> ORGANISM: Homo sapiens

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178 ccccgctgga tcaacgtgct ccaggaggac tctgtgactc tgacatgcca gggggctcgc      180
180 agccctgaga gcgactccat tcagtgggtc cacaatggga atctcattcc caccacacag      240
182 cagcccagct acaggttcaa ggccaacaac aatgacagcg gggagtacac gtgccagact      300
184 ggccagacca gcctcagcga ccctgtgcat ctgactgtgc ttccgaatg gctgggtgctc      360
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202 ggctacatga ctctgaacct cagggcacct actgacgatg ataaaaacat ctacctgact      900
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243 gttgctgctg tagtggcctt gatctactgc aggaaaaagc ggatttcagc caatcccact      780
245 aatcctgacg aggtgacaa agttggggct gagaacacaa tcacctatc acttctcatg      840
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252 <212> TYPE: DNA
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requires explanation for N
see error summary sheet item 9

RAW SEQUENCE LISTING

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TIME: 09:37:07

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Output Set: N:\CRF3\05102002\J027736.raw

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266 gctggggacac ctgcagctcc cccaaaggct gtgctgaaac tcgagcccca gtggatcaac      180
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270 tccattcagt ggttcacaaa tgggaatctc attcccaccc acacgcagcc cagctacagg      300
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307 gaagatctcc caaaggctgt ggtgttctct gagcctcaat ggtacagggt gctcgagaag      120
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327 gtggacacag gactatatatt ctctatgaag aaaagcattc caagctcaac aagggactgg      720
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RAW SEQUENCE LISTING

DATE: 05/13/2002

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TIME: 09:37:07

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Output Set: N:\CRF3\05102002\J027736.raw

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364 tcattctttc cacctgggta ccaagtctct ttctgcttgg tgatgggtact cttttttgca      660
366 gtggacacag gactatatatt ctctgtgaag acaaacattc gaagctcaac aagagactgg      720
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379 <223> OTHER INFORMATION: FcgammaRI      382 <400> SEQUENCE: 9
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389          20          25          30
392 Val Phe Gln Glu Glu Thr Val Thr Leu Gln Cys Glu Val Pro Arg Leu
393          35          40          45
396 Pro Gly Ser Ser Ser Thr Gln Trp Phe Leu Asn Gly Thr Ala Thr Gln
397          50          55          60
400 Thr Ser Thr Pro Ser Tyr Arg Ile Thr Ser Ala Ser Val Lys Asp Ser
401 65          70          75          80
404 Gly Glu Tyr Arg Cys Gln Arg Gly Pro Ser Gly Arg Ser Asp Pro Ile
405          85          90          95
408 Gln Leu Glu Ile His Arg Asp Trp Leu Leu Gln Val Ser Ser Arg
409          100         105         110
412 Val Phe Thr Glu Gly Glu Pro Leu Ala Leu Arg Cys His Ala Trp Lys
413          115         120         125
416 Asp Lys Leu Val Tyr Asn Val Leu Tyr Tyr Gln Asn Gly Lys Ala Phe
417          130         135         140
420 Lys Phe Phe Tyr Arg Asn Ser Gln Leu Thr Ile Leu Lys Thr Asn Ile
421 145         150         155         160
424 Ser His Asn Gly Ala Tyr His Cys Ser Gly Met Gly Lys His Arg Tyr
425          165         170         175
428 Thr Ser Ala Gly Val Ser Val Thr Val Lys Glu Leu Phe Pro Ala Pro
429          180         185         190
432 Val Leu Asn Ala Ser Val Thr Ser Pro Leu Leu Glu Gly Asn Leu Val

```

RAW SEQUENCE LISTING ERROR SUMMARY
PATENT APPLICATION: US/10/027,736

DATE: 05/13/2002
TIME: 09:37:08

Input Set : A:\Sequence List.txt
Output Set: N:\CRF3\05102002\J027736.raw

Please Note:

Use of n and/or Xaa have been detected in the Sequence Listing. Please review the Sequence Listing to ensure that a corresponding explanation is presented in the <220> to <223> fields of each sequence which presents at least one n or Xaa.

Seq#:5; N Pos. 879

VERIFICATION SUMMARY

DATE: 05/13/2002

PATENT APPLICATION: US/10/027,736

TIME: 09:37:08

Input Set : A:\Sequence List.txt

Output Set: N:\CRF3\05102002\J027736.raw

L:11 M:271 C: Current Filing Date differs, Replaced Current Filing Date

L:247 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:5 after pos.:840